

Riboproteins P0, P1, and P2

dsDNA

Nucleosome

Ku

Centromere A

Centromere B

Scl-70

Pm-Scl

RNA-Polymerases 1, 2, and 3

Th

Jo-1

Mi-2

PL7

PL12

SRP

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**IN THE CLAIMS:**

Cancel claim 11 and amend claims 1, 2, 3, 7, and 8 to read as shown in the following clean versions. These claims are reproduced in Appendix B hereto with markings to show the changes made by this amendment.

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- Sub B1  
Ad
- 1                    1. (amended) A method for the identification of a systemic autoimmune  
2 disease in a test subject suspected of suffering from an otherwise unidentified systemic  
3 autoimmune disease selected from the group consisting of systemic lupus erythematosus,  
4 scleroderma, Sjögren's syndrome, polymyositis, dermatomyositis, CREST, and mixed  
5 connective tissue disease, said method comprising:  
6                    (a) analyzing a single biological sample from said test subject for the  
7 presence and amounts of a plurality of autoantibodies to produce a test data set;

8 (b) comparing said test data set to a library of reference data sets, each  
9 reference data set obtained from a biological sample of a reference subject known  
10 to have a systemic autoimmune disease of known identity; and

11 (c) applying pattern recognition means selected from the group consisting  
12 of k-nearest neighbor analysis, multi-linear regression analysis, Bayesian  
13 probabilistic reasoning, and principal component analysis to produce a  
14 statistically derived decision indicating which systemic autoimmune disease said  
15 test subject is suffering from.

1 2. (amended) A method in accordance with claim 1 in which said test  
2 subject is suffering from two otherwise unidentified systemic autoimmune diseases, and  
3 step (c) comprises applying pattern recognition means to produce a statistically derived  
4 decision indicating which two systemic autoimmune diseases said test subject is suffering  
5 from.

1 3. (amended) A method in accordance with claim 1 in which said pattern  
2 recognition means is a member selected from the group consisting of k-nearest neighbor  
3 analysis, multi-linear regression analysis, and Bayesian probabilistic reasoning.

1 7. (amended) A method in accordance with claim 1 in which said  
2 plurality of autoantibodies comprises antibodies to at least fifteen of the following  
3 antigens:

4 SSA 60,  
5 SSA 52,  
6 SSB 48,  
7 Sm BB',  
8 Sm D1,  
9 RNP 68,  
10 RNP A,  
11 RNP C,

12 Fibrillarin,  
13 Riboproteins P0, P1, and P2,  
14 dsDNA,  
15 Nucleosome,  
16 Ku,  
17 Centromere A,  
18 Centromere B,  
19 Scl-70,  
20 Pm-Scl,  
21 RNA-Polymerases 1, 2, and 3,  
22 Th,  
23 Jo-1,  
24 Mi-2,  
25 PL7,  
26 PL12, and  
27 SRP.

1 8. (amended) A method in accordance with claim 1 in which said  
2 plurality of autoantibodies comprises antibodies to each of the following antigens:

3 SSA 60,  
4 SSA 52,  
5 SSB 48,  
6 Sm BB',  
7 Sm D1,  
8 RNP 68,  
9 RNP A,  
10 RNP C,  
11 Fibrillarin,  
12 Riboproteins P0, P1, and P2,

A3